

SEQUENCE LISTING

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<110> WALLACH, David
      BOLDIN, Mark
      VARFOLOMEEV, Eugene
      METT, Igor

<120> MODULATORS OF THE FUNCTION OF FAS/APO1 RECEPTORS

<130> WALLACH=16B

<150> 08/860,082
<151> 1997-08-19

<150> PCT/US95/16542
<151> 1995-12-14

<150> IL 112022
<151> 1994-12-15

<150> IL 112692
<151> 1995-02-19

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<170> PatentIn version 3.0

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ctg ggc aag cgg cga gac ctg gcc agg gcc agc gag ccg agg aca gag      96
Leu Gly Lys Arg Arg Asp Leu Ala Arg Ala Ser Glu Pro Arg Thr Glu
20                               25                               30

ggc gcg cgg agg gcc ggg ccg cag ccc cgg ccg ctt gca gac ccc gcc      144
Gly Ala Arg Arg Ala Gly Pro Gln Pro Arg Pro Leu Ala Asp Pro Ala
35                               40                               45

atg gac ccg ttc ctg gtg ctg ctg cac tcg gtg tcg tcc agc ctg tcg      192
Met Asp Pro Phe Leu Val Leu Leu His Ser Val Ser Ser Ser Leu Ser
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agc agc gag ctg acc gag ctc aag ttc cta tgc ctc ggg cgc gtg gtc      240
Ser Ser Glu Leu Thr Glu Leu Lys Phe Leu Cys Leu Gly Arg Val Val
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aag cgc aag ctg gag cgc gtg cag agc ggc cta gac ctc ttc tcc atg      288
Lys Arg Lys Leu Glu Arg Val Gln Ser Gly Leu Asp Leu Phe Ser Met
85                               90                               95

ctg ctg gag cag aac gac ctg gag ccc ggg cac acc gag ctc ctg cgc      336
Leu Leu Glu Gln Asn Asp Leu Glu Pro Gly His Thr Glu Leu Leu Arg

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Asp Phe Glu Ala Gly Ala Ala Gly Ala Ala Pro Gly Glu Glu Asp			
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Leu Cys Ala Ala Phe Asn Val Ile Cys Asp Asn Val Gly Lys Asp Trp			
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aga agg ctg gct cgt cag ctc aaa gtc tca gac acc aag atc gac agc			528
Arg Arg Leu Ala Arg Gln Leu Lys Val Ser Asp Thr Lys Ile Asp Ser			
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atc gag gac aga tac ccc cgc aac ctg aca gag cgt gtg cgg gag tca			576
Ile Glu Asp Arg Tyr Pro Arg Asn Leu Thr Glu Arg Val Arg Glu Ser			
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Leu Arg Ile Trp Lys Asn Thr Glu Lys Glu Asn Ala Thr Val Ala His			
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ctg gtg ggg gct ctc agg tcc tgc cag atg aac ctg gtg gct gac ctg			672
Leu Val Gly Ala Leu Arg Ser Cys Gln Met Asn Leu Val Ala Asp Leu			
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Val Gln Glu Val Gln Gln Ala Arg Asp Leu Gln Asn Arg Ser Gly Ala			
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Met Ser Pro Met Ser Trp Asn Ser Asp Ala Ser Thr Ser Glu Ala Ser			
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Gly Ala Arg Arg Ala Gly Pro Gln Pro Arg Pro Leu Ala Asp Pro Ala
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Met Asp Pro Phe Leu Val Leu Leu His Ser Val Ser Ser Ser Leu Ser
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Ser Ser Glu Leu Thr Glu Leu Lys Phe Leu Cys Leu Gly Arg Val Val
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Lys Arg Lys Leu Glu Arg Val Gln Ser Gly Leu Asp Leu Phe Ser Met
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Glu Leu Leu Ala Ser Leu Arg Arg His Asp Leu Leu Arg Arg Val Asp
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Asp Phe Glu Ala Gly Ala Ala Ala Gly Ala Ala Pro Gly Glu Glu Asp
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Leu Cys Ala Ala Phe Asn Val Ile Cys Asp Asn Val Gly Lys Asp Trp
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Arg Arg Leu Ala Arg Gln Leu Lys Val Ser Asp Thr Lys Ile Asp Ser
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Ile Glu Asp Arg Tyr Pro Arg Asn Leu Thr Glu Arg Val Arg Glu Ser
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Leu Arg Ile Trp Lys Asn Thr Glu Lys Glu Asn Ala Thr Val Ala His
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Val Gln Glu Val Gln Gln Ala Arg Asp Leu Gln Asn Arg Ser Gly Ala
 225 230 235 240

Met Ser Pro Met Ser Trp Asn Ser Asp Ala Ser Thr Ser Glu Ala Ser
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